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WHAT IS CLAIMED IS:

- 1. An isolated nucleic acid molecule comprising a polynucleotide having a nucleotide sequence at least 96.7% identical to a sequence selected from the group consisting of:
- (a) a polynucleotide fragment of SEQ ID NO:3 or a polynucleotide fragment of the cDNA sequence included in ATCC Deposit No: PTA-2966, which is hybridizable to SEQ ID NO3;
- (b) a polynucleotide encoding a polypeptide fragment of SEQ ID NO:4 or a polypeptide fragment encoded by the cDNA sequence included in ATCC Deposit No: PTA-2966, which is hybridizable to SEQ ID NO:3;
- (c) a polynucleotide encoding a polypeptide domain of SEQ ID NO:4 or a polypeptide domain encoded by the cDNA sequence included in ATCC Deposit No: PTA-2966, which is hybridizable to SEQ ID NO:3;
- (d) a polynucleotide encoding a polypeptide epitope of SEQ ID NO:4 or a polypeptide epitope encoded by the cDNA sequence included in ATCC Deposit No: PTA-2966, which is hybridizable to SEQ ID NO:3;
- (e) a polynucleotide encoding a polypeptide of SEQ ID NO:4 or the cDNA sequence included in ATCC Deposit No: PTA-2966, which is hybridizable to SEQ ID NO:3, having glycine receptor activity;
 - (f) a polynucleotide which is a variant of SEO ID NO:3;
 - (g) a polynucleotide which is an allelic variant of SEQ ID NO:3;
- (h) an isolated polynucleotide comprising nucleotides 4 to 1293 of SEQ ID NO:3, wherein said nucleotides encode a polypeptide corresponding to amino acids 2 to 431 of SEQ ID NO:4 minus the start codon;
- (i) an isolated polynucleotide comprising nucleotides 1 to 1293 of SEQ ID NO:3, wherein said nucleotides encode a polypeptide corresponding to amino acids 1 to 431 of SEQ ID NO:4 including the start codon;
 - (j) a polynucleotide which represents the complimentary sequence (antisense) of SEQ ID NO:3; and
- (k) a polynucleotide capable of hybridizing under stringent conditions to any one of the polynucleotides specified in (a)-(j), wherein said polynucleotide

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does not hybridize under stringent conditions to a nucleic acid molecule having a nucleotide sequence of only A residues or of only T residues.

- 2. The isolated nucleic acid molecule of claim 1, wherein the polynucleotide fragment comprises a nucleotide sequence encoding a human glycine receptor protein.
- 3. A recombinant vector comprising the isolated nucleic acid molecule of claim 1.
 - 4. A recombinant host cell comprising the vector sequences of claim 3.
- 5. An isolated polypeptide comprising an amino acid sequence at least 96.2% identical to a sequence selected from the group consisting of:
- (a) a polypeptide fragment of SEQ ID NO:4 or the encoded sequence included in ATCC Deposit No: PTA-2966;
- (b) a polypeptide fragment of SEQ 1D NO:4 or the encoded sequence included in ATCC Deposit No: PTA-2966, having glycine receptor activity;
- (c) a polypeptide domain of SEQ ID NO:4 or the encoded sequence included in ATCC Deposit No: PTA-2966;
- (d) a polypeptide epitope of SEQ ID NO:4 or the encoded sequence included in ATCC Deposit No: PTA-2966;
- (e) a full length protein of SEQ ID NO:4 or the encoded sequence included in ATCC Deposit No: PTA-2966;
 - (f) a variant of SEQ ID NO:4;
 - (g) an allelic variant of SEQ ID NO:4;
 - (h) a species homologue of SEQ ID NO:4;
- (i) a polypeptide comprising amino acids 2 to 431 of SEQ ID NO:4, wherein said amino acids 2 to 431 comprise a polypeptide of SEQ ID NO:4 minus the start methionine;
 - (j) a polypeptide comprising amino acids 1 to 431 of SEQ ID NO:4;
 - (k) a polypeptide encoded by the cDNA contained in ATCC Deposit No. PTA-2966; and
 - (l) a polypeptide comprising the polypeptide sequence of SEQ ID NO:74;

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- 5 6. The isolated polypeptide of claim 5, wherein the full length protein comprises sequential amino acid deletions from either the C-terminus or the N-terminus.
 - 7. An isolated antibody that binds specifically to the isolated polypeptide of claim 5.
 - 8. A recombinant host cell that expresses the isolated polypeptide of claim 5.
 - 9. A method of making an isolated polypeptide comprising:
 - (a) culturing the recombinant host cell of claim 8 under conditions such that said polypeptide is expressed; and
 - (b) recovering said polypeptide.
 - 10. The polypeptide produced by claim 9.
 - 11. A method for preventing, treating, or ameliorating a medical condition, comprising the step of administering to a mammalian subject a therapeutically effective amount of the polypeptide of claim 5 or the polynucleotide of claim 1.
 - 12. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject comprising:
 - (a) determining the presence or absence of a mutation in the polynucleotide of claim 1; and
 - (b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or absence of said mutation.
 - 13. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject comprising:
 - (a) determining the presence or amount of expression of the polypeptide of claim 5 in a biological sample; and
 - (b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or amount of expression of the polypeptide.
 - 14. An isolated nucleic acid molecule consisting of a polynucleotide having a nucleotide sequence selected from the group consisting of:
- 35 (a) a polynucleotide encoding a polypeptide of SEQ ID NO:2;

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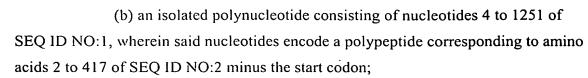
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- (c) an isolated polynucleotide consisting of nucleotides 1 to 1251 of SEQ ID NO:1, wherein said nucleotides encode a polypeptide corresponding to amino acids 1 to 417 of SEQ ID NO:2 including the start codon;
- (d) a polynucleotide which represents the complimentary sequence (antisense) of SEQ ID NO:2;
 - (e) a polynucleotide encoding a polypeptide of SEQ ID NO:4;
- (f) an isolated polynucleotide consisting of nucleotides 4 to 1293 of SEQ ID NO:3, wherein said nucleotides encode a polypeptide corresponding to amino acids 2 to 431 of SEQ ID NO:4 minus the start codon;
 - (g) an isolated polynucleotide consisting of nucleotides 1 to 1293 of SEQ ID NO:3, wherein said nucleotides encode a polypeptide corresponding to amino acids 1 to 431 of SEQ ID NO:2 including the start codon;
 - (h) a polynucleotide encoding the HGRAsv polypeptide encoded by the cDNA clone contained in ATCC Deposit No. PTA-2966; and
 - (i) a polynucleotide which represents the complimentary sequence (antisense) of SEQ ID NO:4.
 - 15. The isolated nucleic acid molecule of claim 14, wherein the polynucleotide comprises a nucleotide sequence encoding a human glycine receptor protein.
 - 16. A recombinant vector comprising the isolated nucleic acid molecule of claim 14.
 - 17. A recombinant host cell comprising the recombinant vector of claim 16.
 - 18. An isolated polypeptide consisting of an amino acid sequence selected from the group consisting of:
 - (a) a polypeptide fragment of SEQ ID NO:2 having glycine receptor activity;
- 35 (b) a polypeptide domain of SEQ ID NO:2 having glycine receptor activity;



a full length protein of SEQ ID NO:2; 5 (c) (d) a polypeptide corresponding to amino acids 2 to 417 of SEQ ID NO:2, wherein said amino acids 2 to 417 comprise a polypeptide of SEQ ID NO:2 minus the start methionine; (e) a polypeptide corresponding to amino acids 1 to 417 of SEQ ID NO:2; a polypeptide fragment of SEO ID NO:4 having glycine receptor 10 (f) activity; (g) a polypeptide domain of SEO ID NO:4 having glycine receptor activity; (h) a full length protein of SEQ ID NO:4; 15 (i) a polypeptide corresponding to amino acids 2 to 431 of SEQ ID NO:4, wherein said amino acids 2 to 431 comprise a polypeptide of SEO ID NO:4 minus the start methionine; (j) a polypeptide corresponding to amino acids 1 to 431 of SEO ID NO:4; and 20 (k) a polypeptide encoded by the cDNA contained in ATCC Deposit No. PTA-2966. 19. The method for preventing, treating, or ameliorating a medical condition of claim 11, wherein the medical condition is a disorder selected from the group consisting of: neural disorder, a neural disorder related to aberrant excitotoxic cell death, a neural disorder 25 related to chronic peripheral neuropathies, a gastrointestinal disorder, a gastrointestinal disorder related to aberrant longitudinal muscle/myenteric plexus contractions, irritable bowel syndrome, a

disorder related to hyper glycine receptor activity.

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